



SEQUENCE LISTING

5 <110> Tillett, D
Thomas, T

<120> A method of sequestering and/or purifying a polypeptide

10 <130> nuc2004

<140> 10/785,452

<141> 2004-02-25

15 <150> PCT/AU02/01159

<151> 2002-08-27

20 <160> 12

<210> 1

<211> 714

25 <212> DNA

<213> Aequorea victoria

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Met Ser Lys Gly Glu Glu Leu Phe Thr Gly Val Val Pro Ile Leu
1 5 10 15

35 gtt gaa tta gat ggc gat gtt aat ggg caa aaa ttc tct gtc agt
90
Val Glu Leu Asp Gly Asp Val Asn Gly Gln Lys Phe Ser Val Ser
20 25 30

40 gga gag ggt gaa ggt gat gca aca tac gga aaa ctt acc ctt aaa
135
Gly Glu Gly Glu Gly Asp Ala Thr Tyr Gly Lys Leu Thr Leu Lys
35 40 45

45 ttt att tgc act act ggg aag cta cct gtt cca tgg cca aca ctt
180
Phe Ile Cys Thr Thr Gly Lys Leu Pro Val Pro Trp Pro Thr Leu
50 55 60

50

50

ccc aac gaa aag aga gat cac atg atc ctt ctt gag ttt gta aca
675
Pro Asn Glu Lys Arg Asp His Met Ile Leu Leu Glu Phe Val Thr
215 220 225

5 gct gct ggg att aca cat ggc atg gat gaa cta tac aaa 714
Ala Ala Gly Ile Thr His Gly Met Asp Glu Leu Tyr Lys
230 235 238

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<211> 1149

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<213> Escherichia coli

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1 5 10 15

25 atc ggc gtc ggc ggc ggc ggc ggt aat gct gtt gaa cac atg gtg
90
Ile Gly Val Gly Gly Gly Gly Gly Asn Ala Val Glu His Met Val
20 25 30

30 cgc gag cgc att gaa ggt gtt gaa ttc ttc gcg gta aat acc gat
135
Arg Glu Arg Ile Glu Gly Val Glu Phe Phe Ala Val Asn Thr Asp
35 40 45

35 gca caa gcg ctg cgt aaa aca gcg gtt gga cag acg att caa atc
180
Ala Gln Ala Leu Arg Lys Thr Ala Val Gly Gln Thr Ile Gln Ile
50 55 60

40 ggt agc ggt atc acc aaa gga ctg ggc gct ggc gct aat cca gaa
225
Gly Ser Gly Ile Thr Lys Gly Leu Gly Ala Gly Ala Asn Pro Glu
65 70 75

45 gtt ggc cgc aat gcg gct gat gag gat cgc gat gca ttg cgt gcg
270
Val Gly Arg Asn Ala Ala Asp Glu Asp Arg Asp Ala Leu Arg Ala
80 85 90

50

50

	gaa atg gct atc tct tct ccg ctg ctg gaa gat atc gac ctg tct	
	765	
	Glu Met Ala Ile Ser Ser Pro Leu Leu Glu Asp Ile Asp Leu Ser	
	245	255
5	ggc gcg cgc ggc gtg ctg gtt aac atc acg gcg ggc ttc gac ctg	
	810	
	Gly Ala Arg Gly Val Leu Val Asn Ile Thr Ala Gly Phe Asp Leu	
	260	270
10	cgt ctg gat gag ttc gaa acg gta ggt aac acc atc cgt gca ttt	
	855	
	Arg Leu Asp Glu Phe Glu Thr Val Gly Asn Thr Ile Arg Ala Phe	
	275	285
15	gct tcc gac aac gcg act gtg gtt atc ggt act tct ctt gac ccg	
	900	
	Ala Ser Asp Asn Ala Thr Val Val Ile Gly Thr Ser Leu Asp Pro	
	290	300
20	gat atg aat gac gag ctg cgc gta acc gtt gtt gcg aca ggt atc	
	945	
	Asp Met Asn Asp Glu Leu Arg Val Thr Val Val Ala Thr Gly Ile	
	305	315
25	ggc atg gac aaa cgt cct gaa atc act ctg gtg acc aat aag cag	
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	Gly Met Asp Lys Arg Pro Glu Ile Thr Leu Val Thr Asn Lys Gln	
	320	330
30	gtt cag cag cca gtg atg gat cgc tac cag cag cat ggg atg gct	
	1035	
	Val Gln Gln Pro Val Met Asp Arg Tyr Gln Gln His Gly Met Ala	
	335	345
35	ccg ctg acc caa gag cag aag ccg gtt gct aaa gtc gtg aat gac	
	1080	
	Pro Leu Thr Gln Glu Gln Lys Pro Val Ala Lys Val Val Asn Asp	
	350	360
40	aat gcg ccg caa act gcg aaa gag ccg gat tat ctg gat atc cca	
	1125	
	Asn Ala Pro Gln Thr Ala Lys Glu Pro Asp Tyr Leu Asp Ile Pro	
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5 <212> DNA

<213> Human rhinovirus

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     1      5      10      15

15 atg act ata aca acc tca aag gga gag ttc aca ggg tta ggc ata
    90
    Met Thr Ile Thr Thr Ser Lys Gly Glu Phe Thr Gly Leu Gly Ile
           20      25      30

20 cat gat cgt gtc tgt gtg ata ccc aca cac gca cag cct ggt gat
   135
    His Asp Arg Val Cys Val Ile Pro Thr His Ala Gln Pro Gly Asp
           35      40      45

25 gat gta cta gtg aat ggt cag aaa att aga gtt aag gat aag tac
   180
    Asp Val Leu Val Asn Gly Gln Lys Ile Arg Val Lys Asp Lys Tyr
           50      55      60

30 aaa tta gta gat cca gag aac att aat cta gag ctt aca gtg ttg
   225
    Lys Leu Val Asp Pro Glu Asn Ile Asn Leu Glu Leu Thr Val Leu
           65      70      75

35 act tta gat aga aat gaa aaa ttc aga gat atc agg gga ttt ata
   270
    Thr Leu Asp Arg Asn Glu Lys Phe Arg Asp Ile Arg Gly Phe Ile
           80      85      90

40 tca gaa gat cta gaa ggt gtg gat gcc act ttg gta gta cat tca
   315
    Ser Glu Asp Leu Glu Gly Val Asp Ala Thr Leu Val Val His Ser
           95      100      105

45 aat aac ttt acc aac act atc tta gaa gtt ggc cct gta aca atg
   360
    Asn Asn Phe Thr Asn Thr Ile Leu Glu Val Gly Pro Val Thr Met
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50 gca gga ctt att aat ttg agt agc acc ccc act aac aga atg att
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cgt	tat	gat	tat	gca	aca	aaa	act	ggg	cag	tgt	gga	ggg	gtg	ctg
450														
Arg	Tyr	Asp	Tyr	Ala	Thr	Lys	Thr	Gly	Gln	Cys	Gly	Gly	Val	Leu
				140					145					150

10

tgt	gct	act	ggg	aag	atc	ttt	ggg	att	cat	gtt	ggc	ggg	aat	gga
495														
Cys	Ala	Thr	Gly	Lys	Ile	Phe	Gly	Ile	His	Val	Gly	Gly	Asn	Gly
				155					160					165

15

aga	caa	gga	ttt	tca	gct	caa	ctt	aaa	aaa	caa	tat	ttt	gta	gag
540														
Arg	Gln	Gly	Phe	Ser	Ala	Gln	Leu	Lys	Lys	Gln	Tyr	Phe	Val	Glu
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aaa	caa	546												
Lys	Gln													
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<223> Polymerase chain reaction oligonucleotide primer

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20 <223> OTHER INFORMATION: Polymerase chain reaction
oligonucleotide primer

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